SEQUENCE LISTING

<110> OOKURA, TETUYA

KASUMI, TAKAFUMI

ASABA, EIJI

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		ttg Leu 195														624
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		cct Pro														720
		cat His														768
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Lys Val Tyr Gln Asn Gln Ser Glu Ile Gly Val Ala Leu Gln Glu Leu 50 55 60

Phe Asp Gln Gly Ile Val Lys Arg Glu Asp Leu Phe Ile Thr Ser Lys 70 75 80

Val Trp Asn Asn Arg His Ala Pro Glu His Val Glu Pro Ala Leu Asp 85 90 95

Glu Thr Leu Lys Glu Leu Gly Leu Ser Tyr Leu Asp Leu Tyr Leu Ile 100 105 110

His Trp Pro Val Ala Phe Lys Phe Thr Thr Pro Gln Glu Leu Phe Pro 115 120 125

Thr Glu Pro Asp Asn Lys Glu Leu Ala Ala Ile Asp Asp Ser Ile Lys 130 135 140

Leu Val Asp Thr Trp Lys Ala Val Val Ala Leu Lys Lys Thr Gly Lys 145 150 155 160

Thr Lys Ser Val Gly Val Ser Asn Phe Thr Thr Asp Leu Val Asp Leu 165 170 175

Val Glu Lys Ala Ser Gly Glu Arg Pro Ala Val Asn Gln Ile Glu Ala 180 185 190

His Pro Leu Leu Gln Gln Asp Glu Leu Val Ala His His Lys Ser Lys 195 200 205

Asn Ile Val Ile Thr Ala Tyr Ser Pro Leu Gly Asn Asn Val Ser Gly 210 220

Lys Pro Pro Leu Thr Gln Asn Pro Gly Ile Glu Ala Thr Ala Lys Arg 225 230 235 240

Leu Asn His Thr Pro Ala Ala Val Leu Leu Ala Trp Gly Ile Gln Arg 245 250 255

Gly Tyr Ser Val Leu Val Lys Ser Val Thr Pro Ser Arg Ile Glu Ser 260 265 270

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gag act ct Glu Thr Le					336
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gct gac cct Ala Asp Pro 130					432
ttg tcc gad Leu Ser Asy 145	Thr Trp L				480
acc aag tog Thr Lys Ser					528

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	ccc Pro															624
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	tac Tyr															816
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- Gly Val Lys Leu Ala Val Lys Ala Gly Tyr Arg His Leu Asp Leu Ala 35 40 45
- Lys Val Tyr Gln Asn Gln Thr Glu Ile Gly Gln Ala Leu Lys Glu Leu 50 55 60
- Phe Asp Glu Gly Val Val Lys Arg Glu Asp Leu Phe Ile Thr Ser Lys 70 75 80
- Leu Trp Asn Asn Arg His Ala Pro Glu His Val Glu Pro Ala Leu Asp 85 90 95
- Glu Thr Leu Lys Glu Leu Gly Leu Ser Tyr Leu Asp Leu Tyr Leu Ile 100 105 110
- His Trp Pro Val Ala Phe Lys Phe Thr Thr Pro Asp Glu Leu Leu Pro 115 120 125
- Ala Asp Pro Thr Asn Lys Asp Leu Ala Tyr Ile Asp Asp Ser Val Lys 130 135 140
- Leu Ser Asp Thr Trp Lys Ala Val Val Ala Leu Lys Lys Thr Gly Lys 145 150 155 160
- Thr Lys Ser Val Gly Val Ser Asn Phe Ser Thr Arg Leu Val Asp Leu 165 170 175
- Val Glu Glu Ala Ser Gly Glu Arg Pro Ala Val Asn Gln Ile Glu Ala 180 185 190

His Pro Leu Leu Gln Gln Asp Glu Leu Val Ala His His Lys Ser Lys 195 200 205

Asn Ile Val Ile Thr Ala Tyr Ser Pro Leu Gly Asn Asn Val Ala Gly 210 215 220

Lys Pro Pro Leu Thr Glu Asn Pro Gly Ile Val Asp Ala Ala Lys Arg 225 230 235 240

Leu Asn His Thr Pro Ala Ala Val Leu Ile Ala Trp Gly Ile Gln Arg 245 250 255

Gly Tyr Ser Val Leu Val Lys Ser Val Thr Pro Ser Arg Ile Lys Ser 260 265 270

Asn Phe Glu Gln Ile Thr Leu Ser Asp Glu Glu Phe Gln Arg Val Thr 275 280 285

Asn Leu Ile Lys Glu Tyr Gly Glu Ser Arg Asn Asn Val Pro Phe Asn 290 295 300

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gat Asp											240
tgg Trp											288
act Thr											336
tgg Trp											384
gac Asp 130											432
tcc Ser											480
aag Lys											528
gag Glu											576

cac His	ccc Pro	ttg Leu 195	ttg Leu	caa Gln	caa Gln	gac Asp	gag Glu 200	ttg Leu	gtt Val	gct Ala	cac His	cat His 205	aag Lys	agc Ser	aag Lys	624
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Asn Ile Val Ile Thr Ala Tyr Ser Pro Leu Gly Asn Asn Val Ala Gly 210 215 220

Lys Pro Pro Leu Thr Glu Asn Pro Gly Ile Val Asp Ala Ala Lys Arg 225 230 235 240

Leu Asn His Thr Pro Ala Ala Val Leu Ile Ala Trp Gly Ile Gln Arg 245 250 255

Gly Tyr Ser Val Leu Val Lys Ser Val Thr Pro Ser Arg Ile Lys Ser 260 265 270

Asn Phe Glu Gln Ile Thr Leu Ser Asp Glu Glu Phe Gln Arg Val Thr 275 280 285

Asn Leu Ile Lys Glu Tyr Gly Glu Ser Arg Asn Asn Val Pro Phe Asn 290 295 300

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